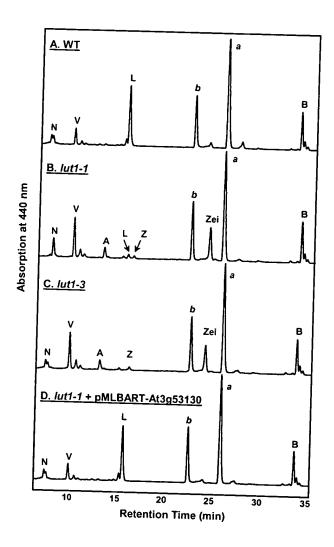
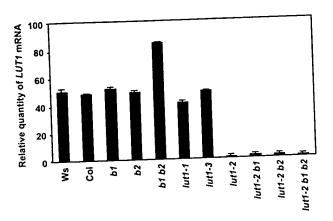
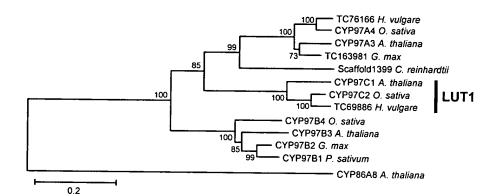


MESSLFSPSSSYSSLFTAKPTRLLSPKPKFTFSIRSSIEKPKPKLETNSS
KSQSWVSPDWLTTLTRTLSSGKNDESGIPIANAKLDDVADLLGGALFLPLY
KWMNEYGPIYRLAAGPRNFVIVSDPAIAKHVLRNYPKYAKGLVAEVSEFLF
GSGFAIAEGPLWTARRAVVPSLHRRYLSVIVERVFCKCAERLVEKLQPYA
EDGSAVNMEAKFSOMTLDVIGLSLFNYNFDSLTTDSPVIEAVYTALKEAEL
RSTDLLPYWKIDALCKIVPRQVKAEKAVTLIRETVEDLIAKCKEIVEREGE
RINDEEYVNDADPSILRFLLASREEVSSVQLRDDLLSMLVAGHETTGSVLT
WTLYLLSKNSSALRKAQEEVDRVLEGRNPAFEDIKELKYITRCINESMRLY
PHPPVLIRRAQVPDILPGNYKVNTGQDIMISVYNIHRSSEVWEKAEEFLPE
RFDIDGAIPNETNTDFKFIPFSGGPRKCVGDQPALMEAIVALAVFLQRLNV
ELVPDQTISMTTGATIHTTNGLYMKVSQR







Dissociation energy: 100 kcal/mol

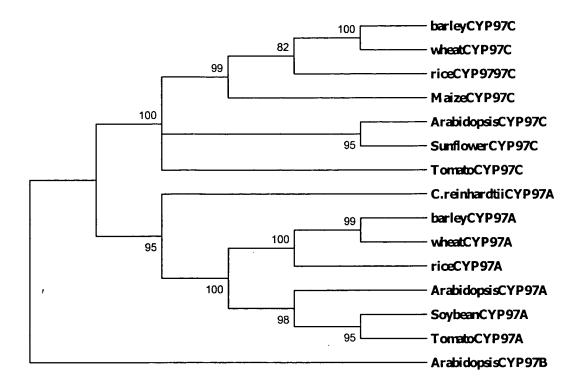
Dissociation energy: 86 kcal/mol

Figure 7

CYP97A3(Arabidopsis) single knockout mutant (SALK_116660)



Figure 8



Phylogenetic tree (Neighbor-joining tree with p-distance, Pairwise deletion method was used. Arabidopsis CYP97B is an outgroup.)

Amino acid similarity

	CYP97A	CYP97C
Arabidopsis	100%	100%
Rice	405/544 (74%).	385/488 (78%)
Barley	374/481 (77%).	395/524 (75%)
G. max(Soybean)	343/410 (83%)	not included
Wheat	243/328 (74%)	264/2E0 /720/)
tomato	(0/1/) (10/0/)	234/330 (72%),
	44 1/346 (80%),	226/279 (81%),
suni Iower	not included	167/202 (82%),
maize	not included	145/177 (81%).
Chlamydomonas	223/365 (61%),	not included

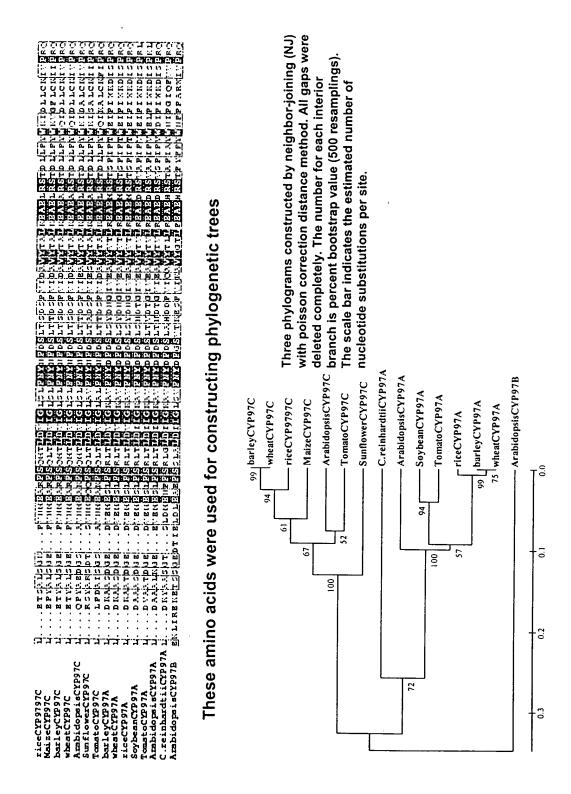
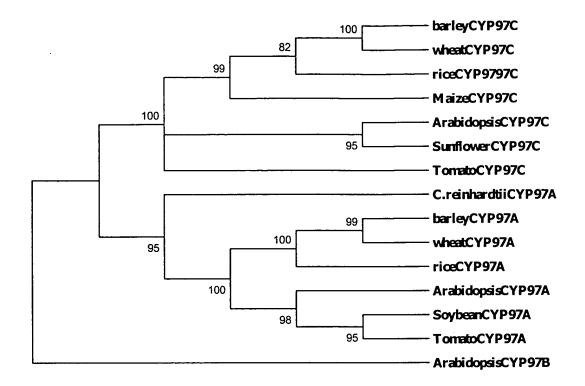
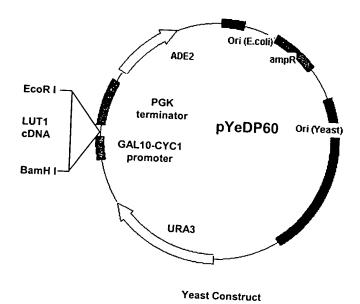


Figure 11



Phylogenetic tree

(Neighbor-joining tree with p-distance, Pairwise deletion method was used. Arabidopsis CYP97B is an outgroup.)



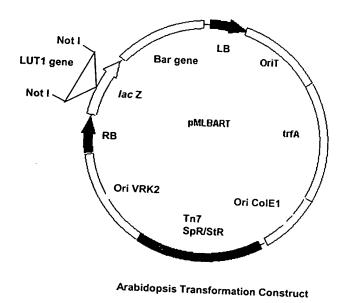


Table 1. $\beta\textsc{-Xanthophyll}$ production and $\beta\textsc{-ring}$ hydroxylation in leaf tissue of wild type and carotenoid hydroxylase mutants $\dot{}$

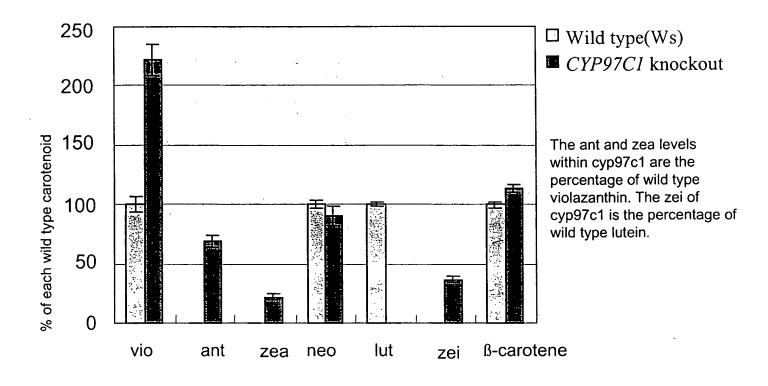
Genotype	β-Xanthophylls [†]	Hydroxylated β-rings [‡] 48.5 ± 1.0 ^a			
Ws	54.0 ± 2.7°§				
Col	60.7 ± 7.6^{a}	48.7 ± 0.9^{a}			
b1 b2	20.5 ± 4.8^{b}	40.2 ± 1.4 ^b			
lut1-2 b1 b2	26.5 ± 3.4^{b}	$33.6 \pm 2.4^{\circ}$			
lut1-3 b1 b2	28.3 ± 4.6^{b}	31.1 ± 1.2°			

Total carotenoids were extracted from five-week-old plants and quantified by HPLC as previously described (Tian et al., 2003). † β -xanthophylls are the sum of zeaxanthin, antheraxanthin, violaxanthin, and neoxanthin as mmol pigment/ mol chlorophyll a+b.

[‡] Data are given as percentage of total ring hydroxylation.

[§] All values are means ± SD (n = 6). Values marked with the same letters are not significantly different from each other within a column (Student's t test, P > 0.05).

Figure 14



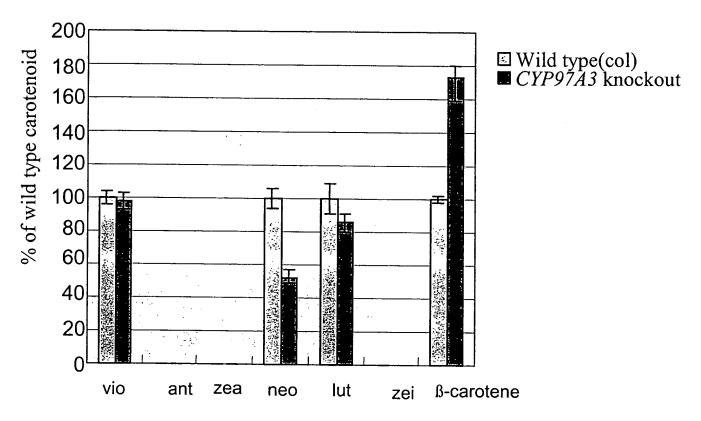
type and CYP97C1 knockout (p>0.05)

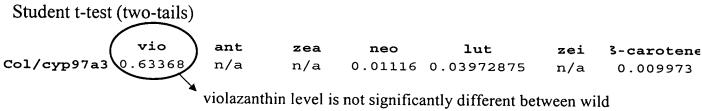
Student t-test (two-tails)

vio ant zea neo lut zei ß- carotene
Ws/cvo97c1 0.007098 0.001631 0.00604 0.111459 9.22469E-05 0.00153 0.04776516

Neoxanthin level is not significantly different between wild

Figure 15





violazanthin level is not significantly different between wild type and CYP97CI knockout (p>0.05)

Figure 16

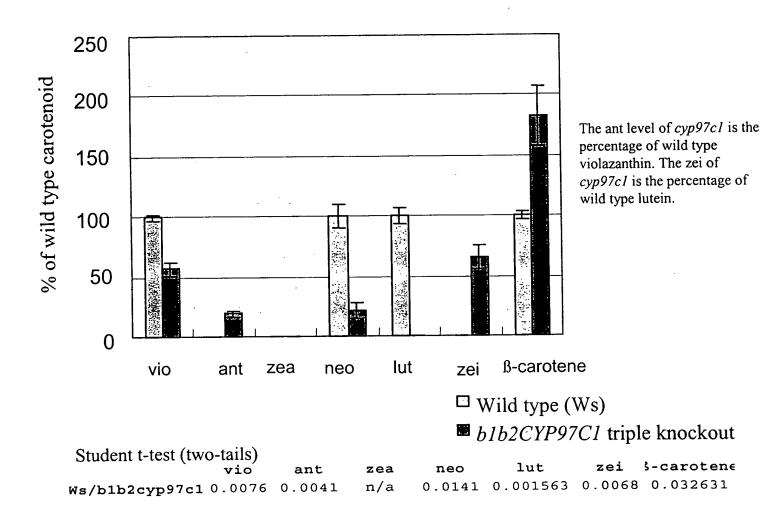


Figure 17

	vio/(chla+b)	ant/(chla+b)	zea/(chla+b)	chla+b)	lut/(chla+b)	rei/(a+b)	car/(chla+b
	vio/(c	ant/(c	zea/(neo/(chla	lut/(zei,	خّ
wild type(Ws)	100	0	0	100	100	0	100
STDEV, wild type(Ws)	6.1	0	0	3.17	1.66	0	2.43
CYP97C1 knockout	221.8388	69.77	21.94	89.97	0	36.82	113.17
STDEV, CYP97C1 knockout	12.72	4.89	2.97	8.14	0	2.49	2.96
wild type(col)	100	0	0	100	100	0	100
STDEV, wild type (col)	3.78	0	0	5.83	8.67	0	
CYP97A3 knockout	98.12	0	0	52	86.2	0	173.49
STDEV, CYP97A3 knockout (col)	5.05	0	0	3.03	6.13	0	14.42
wild type(Ws)	100	0	0	100	100		
STDEV, wild type(Ws)	2.36	0	0	9.83	6.86	0	
b1b2CYP97C1 triple knockout	57.15	19.36	0	21.17	0	65.62	182.97
STDEV, b1b2CYP97C1 triple knock	5.39	2.14	0	7.16	0	9.43	24.7

Fig. 18

SEQ ID NO: 1: CYP97C Arabidopsis thaliana:
LQPYAEDGSAVNMEAKFSQMTLDVIGLSLFNYNFDSLTTDSPVIEAVYTALKEAELRSTDLLPYWKIDALC
KIVPRQ

SEQ ID NO: 2: CYP97A Arabidopsis thaliana:
LDAAAI,KGEEVEMESLFSRLTLDIIGKAVFNYDFDSLTNDTGVIEAVYTVLREAEDRSVSPIPVWDIPIWK
DISPRQ

SEQ ID NO: 3: CYP97B Arabidopsis thaliana: EKLIREKETSSGEDTIELDLEAEFSSLALDIIGLSVFNYDFGSVTKESPVIKAVYGTLFEAEHRSTFYFPY WNFPPARWIVPRQ

Figure 19a

SEQ ID NO: 4: LUT1 Arabidopsis thaliana (CYP97C1):

MESSLFSPSSSYSSLFTAKPTRLLSPKPKFTFSIRSSIEKPKPKLETNSSKSQSWVSPDWLTTLTRTLSS
GKNDESGIPIANAKLDDVADLLGGALFLPLYKWMNEYGPIYRLAAGPRNFVIVSDPAIAKHVLRNYPKYAK
GLVAEVSEFLFGSGFAIAEGPLWTARRAVVPSLHRRYLSVIVERVFCKCAERLVEKLQPYAEDGSAVNME
AKFSQMTLDVIGLSLFNYNFDSLTTDSPVIEAVYTALKEAELRSTDLLPYWKIDALCKIVPRQVKAEKAVT
LIRETVEDLIAKCKEIVEREGERINDEEYVNDADPSILRFLLASREEVSSVQLRDDLLSMLVAGHETTGSV
LTWTLYLLSKNSSALRKAQEEVDRVLEGRNPAFEDIKELKYITRCINESMRLYPHPPVLIRRAQVPDILPG
NYKVNTGQDIMISVYNIHRSSEVWEKAEEFLPERFDIDGAIPNETNTDFKFIPFSGGPRKCVGDQFALMEA
IVALAVFLQRLNVELVPDQTISMTTGATIHTTNGLYMKVSQR

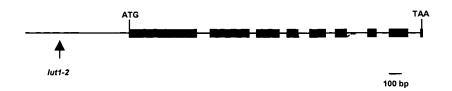
SEQ ID NO: 5: LUT1 Arabidopsis thaliana (CYP97C1): tttatcaccaaaacccaaattcacattctccatcagatcctccattgagaaacccaaacccaaactcgaga ggaaaaaacgacgagtcaggtataccaatcgcgaacgcgaagctcgacgatgtcgctgatctcctcqqaqg tqctcttcttcttacctctctacaaatqqatqaatqaqtacqqacccatttaccqtctcqctqctgqtcctc qtaatttcqtaattqtqaqcqacccaqcqataqctaaacatqttttqaqqaattatccaaagtacqctaaa qqcttaqtcqctqaaqtctctqaatttctatttqqttcqqqtttcqctatcqctqaaqqacctctttggac agtaatttcatctcctcctatctcaattttgaagtttttggaattgtgggaagtaatgtgtgactgtcttgt atgataagtaactctaattttagggtttagattccaatcttctctattgggcttagctgaagtctgatttt ttacataqqcqaqqcqtaqaqcqqtqqttccatcqcttcacaqqaqqtatttqtctgtgattgtggagaga gtattctgcaaatgtgcagagaggcttgttgagaagttgcagccttatgcagaagacggaagtgctgtgaa tatggaagcgaagttctctcagatgacacttgatgtcattgggttgtctctttttaactacaatttcgatt ctttqactactqataqtcctqtcattqaaqctqtttacactqctcttaaaqaaqctqaqcttcqttctact gatcttctgccatattggaaggcaagtttcctgtgtttttttctgtggtttgttgattgtgtggaacaattg gattettgttaattgagagggtttggttgtttttttcagategatgeattgtgtaagatagteeegagaea ggtgaaagctgaaaaggctgtaactttgataagggaaactgttgaagaccttattgctaagtgtaaagaaa ttgtcgaaagagaaggcgaaagaatcaatgatgaggagtatgtaaatgatgctgacccaagtatcctgcgt ttcttgcttgcaagcagagaagaggtttaaacttttttccttaagtttataagcaaatttggcctttcatt atcgcataatcgaagctgatgttgcattgtgagggttttcaggtatcaagtgtgcagttacgggatgatct teteteaatgetegtagegggteatgaaaceaetggatetgteeteaettggaeaetttateteetaagta aggtaccttaatgtatcttctactttgctatgctagagaatttacttggatgggagcttctctgttctcat ttacctcttcaaattctctatgttcatagaactcatctgcattaaggaaagcacaagaagaagtagacaga qtqttaqaaqqaaqaaacccqqctttcqaqqatataaaqqaqttqaaqtacatcactcqttqtataaacqa gtcaatgcgtctctatcctcatcctcctgtaagcaatcaagctcatctctctaattattcatgaactaaat tttctgattgatttgtttcctggtaggtcttgataagaagagctcaagttcctgacattcttcctgggaac ctcttccttctctcgtccatagtataacataggggagcctaatccttctcttcaatgatctttgtgtgtt ttattccaaaggtatgggaaaaagctgaggaatttctgcctgaacgattcgacatagatggcgcaatccct aacqaaacaaacactqatttcaaqtaaactcaqtaqaacacatcttttqacacaaactactqaatcaaqat tagtqqttttqattaqqqaatttaaaaqatqattttcttttttcaccaqattcatcccattcaqtqqaqqq cctagaaaatgtgtaggcgatcagtttgcattgatggaggcaattgtggcactcgcggtgtttcttcagcg gttaaacgttgagctggttcctgatcagaccattagcatgaccacaggagcaaccatacacaccaccaatg tatgccaatgttctcacactcgagagattaatgagagtgtctgttttgtttagaatgattccaatttctct aatgctgatattttcaatttcagggattgtatatgaaggtgagccaaaggtaa

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Figure 20

mutant LUT1-1 Arabidopsis thaliana (lut1-1): SEQ ID NO: 7: tttatcaccaaaacccaaattcacattctccatcagatcctccattgagaaacccaaacccaaactcgaga qqaaaaaaacqacqaqtcaqqtataccaatcqcqaacqcqaaqctcqacqatqtcqctqatctcctcqqaqq tgctctcttcttacctctctacaaatqqatqaatqaqtacqqacccatttaccqttctcqctqctqqtcctc qtaatttcqtaattqtqaqcqacccaqcqataqctaaacatgttttqaqqaattatccaaagtacqctaaa qqcttaqtcqctqaaqtctctqaatttctatttqqttcqggtttcgctatcgctgaaggacctctttggac aqtaatttcatctcctcctatctcaattttqaaqttttttqqaattqtqqaaqtaatqtqtqactgtcttgt atqataaqtaactctaattttaqqqtttaqattccaatcttctctattqqqcttaqctqaaqtctgatttt ttacataggcgaggcgtagagggtggttccatcgcttcacaggaggtatttgtctgtgattgtggagaga gtattctgcaaatgtgcagagggcttgttgagaagttgcagccttatgcagaagacggaagtgctgtgaa tatggaagcgaagttctctcagatgacacttgatgtcattgggttgtctctttttaactacaatttcgatt $\verb|ctttgactactgatagtcctgtcattgaagctgtttacactgctcttaaagaagctgagcttcgttctact|\\$ $\tt gatcttctgccatattggaaggcaagtttcctgtgttttttctgtggtttgttgattgtgtggaacaattg$ qattcttqttaattqaqaqqqtttqqttqttttttttcaqatcgatgcattqtgtaagatagtcccgagaca ggtgaaagctgaaaaggctgtaactttgataagggaaactgttgaagaccttattgctaagtgtaaagaaa ttgtcgaaagagaaggcgaaagaatcaatgatgaggagtatgtaaatgatgctgacccaagtatcctgcgt ttcttgcttgcaagcagagagaggtttaaacttttttccttaagtttataagcaaatttggcctttcatt atcgcata atcgaagctgatgttgcattgtgagggttttcaggtatcaagtgtgcagttacgggatqatcttctctcaatqctcqtaqcqqqtcatqaaaccactqqatctgtcctcacttgqacactttatctcctaagta aqataccttaatgtatcttctactttgctatgctagagaatttacttggatgggagcttctctgttctcat ttacctcttcaaattctctatqttcataqaactcatctqcattaaqqaaaqcacaaqaaqaaqtaqacaqa gtgttagaaggaagaaacccggctttcgaggatataaaggagttgaagtacatcactcgttgtataaacga gtcaatgcgtctctatcctcatcctcctgtaagcaatcaagctcatctctctaattattcatgaactaaat tttctgattgatttgtttcctggtaggtcttgataagaagagctcaagttcctgacattcttcctgggaac ctcttccttctctcgtccatagtataacataggggagcctaatccttctcttcaatgatctttgtgtggtt ttattccaaaggtatgggaaaaagctgaggaatttctgcctgaacgattcgacatagatggcgcaatccct aacgaaacaacactgatttcaagtaaactcagtagaacacatcttttgacacaaactactgaatcaagat tagtggttttgattagggaatttaaaagatgattttctttttttcaccagattcatcccattcagtggaggg cctagaaaatgtgtaggcgatcagtttgcattgatggaggcaattgtgggcactcgcggtgtttcttcagcg gttaaacgttgagctggttcctgatcagaccattagcatgaccacaggagcaaccatacacaccaccaatg tatgccaatgttctcacactcgagagattaatgagagtgtctgttttgtttagaatgattccaatttctct aatgctgatattttcaatttcagggattgtatatgaaggtgagccaaaggtaa

Figure 21
SEQ ID NO: 8: LUT1-2 mutant Arabidopsis thaliana (lut1-2):



SEQ ID NO: 9: mutant Arabidopsis thaliana LUT1-3 (lut1-3)

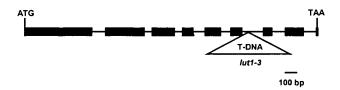


Figure 22

SEQ ID NO: 10: conserved transmembrane domain: LVAEVSEFLFGSGFAIAEGPLWTA

SEQ ID NO: 11: conserved an N-terminal transit peptide for chloroplast-targeting: MESSLFSPSSSSYSSLFTAKPTRLLSPKPKFTFSIR

SEQ ID NO: 12: conserved consensus motif of cytochrome P450 monooxygenase molecular oxygen binding pocket: (A/G)GX(D/E)T(T/S)

SEQ ID NO: 13: conserved sequence of cytochrome P450 monooxygenase molecular oxygen binding pocket in LUT1 Arabidopsis thaliana: AGHETT

SEQ ID NO: 14: conserved consensus cysteine motif in p450 monooxygenase enzymes: FXXGXXXCXG

SEQ ID NO: 15: conserved cysteine sequence in LUT1 Arabidopsis thaliana: FSGGPRKCVG

SEQ ID NO: 16: rice CYP97C2 Oryza sativa:

MAAAAAAAVPCVPFLCPPPPPLVSPRLRRGHVRLRLRPPRSSGGGGGGGGGGDEPPITTSWVSPDWLTALSR
SVATRLGGGDDSGIPVASAKLDDVRDLLGGALFLPLFKWFREEGPVYRLAAGPRDLVVVSDPAVARHVLRGY
GSRYEKGLVAEVSEFLFGSGFAIAEGALWTVRRRSVVPSLHKRFLSVMVDRVFCKCAERLVEKLETSALSGK
PVNMEARFSQMTLDVIGLSLFNYNFDSLTSDSPVIDAVYTALKEAELRSTDLLPYWKIDLLCKIVPRQIKAE
KAVNIIRNTVEDLITKCKKIVDAENEQIEGEEYVNEADPSILRFLLASREEVTSVQLRDDLLSMLVAGHETT
GSVLTWTIYLLSKDPAALRRAQAEVDRVLQGRLPRYEDLKELKYLMRCINESMRLYPHPPVLIRRAIVDDVL
PGNYKIKAGQDIMISVYNIHRSPEVWDRADDFIPERFDLEGPVPNETNTEYRFIPFSGGPRKCVGDQFALLE
AIVALAVVLQKMDIELVPDQKINMTTGATIHTTNGLYMNVSLRKVDREPDFALSGSR

SEQ ID NO: 17: barley CYP97C Hordeum vulgare:
MPAAAFASALASPPPPWAPRPSPRHASLRLPPPRSSGGGGDKPTTSWVSPDWLTSLSRSVLGRGNDDSGIPV
ASAKLDDVQDLLGGALFLPLFKWFREEGPVYRLAAGPRDFVIVSDPAVAKHVLRGYGTRYEKGLVAEVSEFL
FGSGFAIAEGALWTVRRRAVVPSLHKRFLSVMVDKVFCKCAERLVEKLETYALSGEPVNMEARFSQMTLDVI
GLSLFNYNFDSLTSDSPVIDAVYTALKEAEARSTDLLPYWQIDLLCKIVPRQIKAEKAVNTIRNTVEELIIK
CKAIVDAENEQIEGEEYVNEADPSILRFLLASREEVSSLQLRDDLLSMLVAGHETTGSVLTWTIYLLSKDPV
ALRRAQDEVDRVLQGRLPRYEDVKELKYLMRCINESMRLYPHPPVLIRRALVDDVLPGNYKVKTGQDIMISV
YNIHRSPEVWDRADEFIPERFDLEGPIPNESNTDFRFIPFSGGPRKCVGDQFALLEAIVALAIVIQKMDVQL
VADQKISMTTGATIHTTNGLYMNVXLRKVEQEADLALSPSG

SEQ ID NO: 18: wheat CYP97C Triticum aestivum:
MPAAAFASAFASPPPPWAPRPPPRHASLRLPPPRSSSNNSGGGGGDKPTTSWVSPDWLTSLSRSVLGRGNDD
SGIPVASAKLDDVQDLLGGALFLPLFKWFREEGPVYRLAAGPRDFVIVSDPAVAKHVLRGYGTRYEKGLVAE
VSEFLFGSGFAIAEGALWTVRRRAVVPSLHKRFLSVMVDKVFCKCAERLVEKLETYALSGEPVNMEARFSQM
TLDVIGLSLFNYNFDSLTSDSPVIDAVYTALKEAEARSTDLLPYWQIDLLCKIVPRQIKAEKAVNTIRNTVE
ELITKCKAIVDAENEQIEGEEYVNEADPSILRFLLASREEVSSLQLRDDLLSMLVAGHETTGSVPDYRLQAQ

SEQ ID NO: 19: tomato CYP97C Lycopersicon esculentum:
CRCAERMVEKLLPDAISGSAVNMEAKFSQLTLDVIGLALFNYNFDSLTTDSPVIDAVYTALKEAELRSTDLL
PYWQIKALCKFIPRQIKAENAVSLIRQTVEELIAKCREIVETEGERINEDEYVNDRDPSILRFLLASREEVS
SLQLRDDLLSMLVAGHETTGSVLTWTAYLLSKDPSSLEKAHEEVDRVLGGRSPTYEDMKNLKFLTRCITESL
RLYPHPPVLIRRAQVADVLPGNYKVNVGQDIMISVYNIHHSSKVWDRAEEFDPERFDLERSRP

SEQ ID NO: 20: maize CYP97C Zea mays:
LEPYALSGEPVNMEARFSQLTLDVIGLSLFNYNFDSLTTDSPVIDAVYTALKEAELRSTDLLPYWKVGFLCK
IIPRQIKAENAVTIIRNTVEELIMKCKEIVEAENEQIEGEEYVNEGDPSILRFLLASRDEVSSVQLRDDLLS
MLVAGHETTGSVLTWTIYLLSKDPTALRRAQDE

SEQ ID NO: 21: sunflower CYP97C Helianthus annuus:
GPRNFVIVSDPEIAKHVLRNYGSIYAKGLVAEVSEFLFGSGFAIAEGSLWTARRRAVVPSLHKKYLSVIVDR
VFCKCSERLVEKLRSYARSDTSVNMEQQFSQLTLDVIGLAVFNYNFDSLTADSPVIESVYTALKEAEARSTD
LLPYWKISALCKIIPRQIKAEQAVTVIRETVEELIIKCKEIVEKEGEKIDDEDYVNDATYIFIC

Figure 24a

SEQ ID NO: 22: rice CYP97C Oryza sativa:

atggccgccgccgccgccgccgtcccgtgcgtaccattcctgtgcccgcctcctccgccattggtctcg ccgcgtctccgccgtggccacgtccgcctccgcctgcggccgccaaggagcagcggcggtggaggcggaggc ggagcggggggagacgagccgccatcaccacctcgtgggtgagccccgactggctcacggcgctctcccgc tcggtggcaacccgcctcggcggggggcgacgactcggggatcccccgtcgcctccgccaagctcgacgacgtg cgggacctcctcggcggcgctcttcctccctctcttcaagtggttccgcgaggaaggccccgtctaccgc ctcgcggcggggccgcgggatctcgtcgtcagcgatcccgccgttgccaggcacgtgctgcgtgggtac ggttcgaggtacgagaaggggctcgtcgccgaggtttccgagttcctcttcggctccgggttcgccatcgcc gagggcgctctctggacggtgagacgtcgatcagttgtaccatctctacacaaacgatttctctcggtgatg gttgacagagttttttgtaaatgtgctgagagattagtggagaagcttgagacatctgctttaagtggcaaa $\verb|cctgtaaatatggaagcaaggttctctcaaatgactttagatgtgattggtttgtccttgttcaattacaat|\\$ tttgattccctcacatcagatagccctgttattgatgctgtttacactgcactcaaggaagcagaacttcgt tctacagatcttttaccatactggaagattgatttgctgtgcaagattgttcctagacaaataaaagcagaa aaggcagttaacatcatcaggaataccgttgaggacctaattaccaaatgcaagaagattgtagatgctgag aatgaacaaattgagggtgaggaatatgtaaatgaggcagaccctagcatcctgcgattcctacttgctagc cgtgaagaggtaaccagtgtgcagttacgtgatgatctattgtcaatgttagttgctggtcatgaaacaaca ggctctgtactgacgtggactatttatcttctcagtaaggatccagcagcgctgaggagagctcaagcagag gttgaccgtgttctacaaggtagactccccagatatgaagatctaaaagagctgaagtacttgatgcgctgt ataaatgagtctatgcggctttatccacacccacctgtgttgatacggcgagccatagttgatgatgtgctt cccggaaactataagatcaaagctggtcaagatattatgatttcagtgtacaatatacacaggtcacctgag gtttgggacagagctgatgattttattcctgagagatttgatttagagggacctgttccaaatgagacaaac actgaatacagatttatcccattcagtggaggtcctcggaaatgtgttggagatcagtttgctctcttggaa gcaattgtggcacttgctgttgttgcagaagatggacattgagcttgtgccagatcaaaaaattaacatg actactggggccacaattcatacaaccaatggcctgtatatgaatgtaagtctgcgtaaagttgacagggaa $\verb"cctgattttgcactcagtgggtccagatga"$

SEQ ID NO: 23: barley CYP97C Hordeum vulgare:

atgcccgccgcggcattcgcctccgcgctcgcgtctcctcctcctccatgggccccacgaccgtcccctcgg agccccgactggctcacgtcgctgtcccgctcggtgctcggccggggaaacgacgactcggggatccccgtc gcctccgccaagctcgacgtgcaggacctcctcgggggcgcgctcttcctcccgctcttcaagtggttc egegaggaagggeeegtetacegeetegeegggggeegegegacttegteategteagegaeeeegeegtg gccaagcacgtcctccgcgggtacggcacgcggtacgagaaggggctcgtcgccgaggtctccgagttcctc tttggctctgggttcgccatcgccgagggagcgctctggacggtgagacgtagagcagttgtaccatctcta cacaaaagatttctctcagtaatggttgataaagtgttttgtaaatgtgctgagagattggtggaaaagctc gagacatatgctttgagcggtgaacctgttaatatggaagcgagattttctcaaaatgacactagatgtgatt ggtttgtctttgttcaactacaactttgattccctcacatcagatagtcctgttattgatgctgtttacacc gttcctagacagatcaaagcagaaaaggcagttaacacaataaggaatactgttgaagagctaattataaaa tgcaaggcaatcgtagatgctgaaaatgaacagattgagggtgaagaatatgtaaatgaggcagatcctagc ${\tt atcctgcgttttttacttgctagccgtgaagaggtcagcagtttgcagttacgtgatgatctattgtcaatg}$ $\verb|ttagttgctggtcacgaaacaacaggctctgtactgacatggactatttatcttctcagtaaggatccagta|\\$ gcactaaggagagcccaagatgaggtagatcgtgttctacaaggtagactcccaagatatgaagatgtaaaa gagetgaagtaettgatgegetgtateaatgagteeatgeggetataeceaeateeteetgtgetgataegg cgtgcactagttgatgatgtgcttcctggaaactacaaggttaagactggtcaagatattatgatttctgtg ggtcccattccaaatgagtcaaacaccgatttcaggtttatccctttcagtggaggtcctcgaaaatgtgtt ggagatcagtttgctcttttagaagcaattgtggcacttgcaattgtcatacaaaagatggacgttcagctt agnetgegtaaagttgageaagaagetgaettageaetgagteeateaggetag

Figure 24b

SEQ ID NO: 24: wheat CYP97C Triticum aestivum:

atgcccgccgcggcattcgcctccgcgttcgcgtctcctcctcctccgtgggccccacgaccgcctcctcgc cacgccagcctccgcctgcccccgccaaggagcagcaacaacaacagcggcggcggagggggacaagccc teggggataceegtegeeteegeeaagetegaegtgeaggaeeteetegggggegegetetteetgeeg ctcttcaagtggttccgcgaggaagggccgtctaccgcctcgccgcggggccgcgcgacttcgtcatcgtc agcgaccccgccgtagccaagcacgtcctccgcgggtacggcacgcggtacgagaaggggctcgtcgccgag gtctccgagttcctctttggctctgggttcgccatcgccgagggagcgctctggacggtgagacgtagagca gttgtaccatctctacacaaaagatttctctcagtaatggtcgataaagtgttctgtaaatgtgctgagaga ttggtggaaaagctcgagacttatgctttgagtggtgaacctgttaatatggaagcgaggttttctcaaatg acattagatgtgattggtttatccttgttcaactacaactttgattccctcacatcagatagtcctgttatt gatgctgtttacactgcactcaaagaagctgaggctcgttctacagatcttttaccatactggcagatcgat ttgctgtgcaagattgttcctagacagataaaagcggaaaaagcagttaacacaataaggaataccgttgaa gagctaattacaaaatgcaaggcaatcgtagatgctgaaaatgaacagattgagggtgaagaatatgtaaat gaggcagatcctagcatcctgcggtttttacttgctagccgtgaagaggtcagcagtttgcagttacgtgat gatctattgtcaatgttagttgctggtcatgaaacaacaggttctgtaccagactatcgattacaagcccaa ggttcc

Figure 25

SEQ ID NO: 28: forward At3g53130 primer 5'-CTTCCTCTTCTTCTCTCTCTCTCTCTCACT-3'

SEQ ID NO: 29: reverse At3g53130 primer
5'-AAGAACGATGGATGTTATAGACTGAAATC-3'

SEQ ID NO: 30: LUT1 TaqMan probe 5'-CCGTCTCGCTGCTGGTCCTCG-3'

SEQ ID NO: 31: forward LUT1 primer
5'-GGATGAATGAGTACGGACCCAT-3'

SEQ ID NO: 32: reverse *LUT1* primer 5'-GGGTCGCTCACAATTACGAAA-3'

Figure 26a

SEQ ID NO: 33: CYP97A3 Arabidopsis thaliana:

MAMAFPLSYTPTITVKPVTYSRRSNFVVFSSSSNGRDPLEENSVPNGVKSLEKLQEEKRRAELSARIASGA
FTVRKSSFPSTVKNGLSKIGIPSNVLDFMFDWTGSDQDYPKVPEAKGSIQAVRNEAFFIPLYELFLTYGGI
FRLTFGPKSFLIVSDPSIAKHILKDNAKAYSKGILAEILDFVMGKGLIPADGEIWRRRRRAIVPALHQKYV
AAMISLFGEASDRLCQKLDAAALKGEEVEMESLFSRLTLDIIGKAVFNYDFDSLTNDTGVIEAVYTVLREA
EDRSVSPIPVWDIPIWKDISPRQRKVATSLKLINDTLDDLIATCKRMVEEEELQFHEEYMNERDPSILHFL
LASGDDVSSKQLRDDLMTMLIAGHETSAAVLTWTFYLLTTEPSVVAKLQEEVDSVIGDRFPTIQDMKKLKY
TTRVMNESLRLYPQPPVLIRRSIDNDILGEYPIKRGEDIFISVWNLHRSPLHWDDAEKFNPERWPLDGPNP
NETNQNFSYLPFGGGPRKCIGDMFASFENVVAIAMLIRRFNFQIAPGAPPVKMTTGATIHTTEGLKLTVTK
RTKPLDIPSVPILPMDTSRDEVSSALS

SEQ ID NO: 34: rice CYP97A Oryza sativa:

MAATSSAAAAAPPPCRLLGSGQAHLRLPPSAAAAAASARRRLLLRCAASGGNGKGGGGDGSGSDPVLEERRR RRQAELAARIASGEFTAQGPAWIAPLAVGLAKLGPPGELAAALLTKVAGGGGPEIPQAVGSMSAVTGQAFFI PLYDLFLTYGGIFRLNFGPKSFLIVSDPAIAKHILRDNSKAYSKGILAEILEFVMGTGLIPADGEIWRVRRR AIVPAMHQKYVTAMISLFGYASDRLCQKLDKAATDGEDVEMESLFSRLTLDVIGKAVFNYDFDSLSYDNGIV EAVYVTLREAEMRSTSPIPTWEIPIWKDISPRQKKVNEALALINKTLDELIDICKRLVEEEDLQFHEEYMNE QDPSILHFLLASGDDVSSKQLRDDLMTMLIAGHETSAAVLTWTFYLLSKYPNVMAKLQDEADTVLGDRLPTI EDVKKLKYTTRVINESLRLYPQPPVLIRRSIEEDMLGGYPIGRGEDIFISVWNLHHCPKHWDGADVFNPERW PLDGPNPNETNQNFSYLPFGGGPRKCVGDMFATFETVVATAMLVRRFDFQMAPGAPPVEMTTGATIHTTEGL KMTVTRRTKPPVIPNLEMKVISDSPENMSTTTSMPVSAASIASGEDQQGQVSATRI

SEQ ID NO: 35: barley CYP97A Hordeum vulgare:
SARGQAVGSLASVAGEAFFLPLYDLFLTYGGVFRLNFGPKSFLIVSDPDVAKHILRDNSKAYSKGILAEILE
FVMGTGLIPADGEVWRVRRRAIVPALHQKYVTAMIGLFGNASDRLCQKLDKAASDGEDVEMESLFSRLTLDV
IGKAVFNYDFDSLSYDNGIVEAVYVTLREAEMRSTSPIPTWEIPIWKDISPRQRKVNEALALINNILDELIA
TCKRMVDEEDLQFHEEYMNEKDPSILHFLLASGDDVSSKQLRDDLMTMLIAGHETSAAVLTWTFYLLSKYPN
VMSKLQAEADAVLGDGLPTIDDVKKLKYTTRVINESLRLYPQPPVLIRRSLEDDMLGEYPIGKGEDIFISIW
NLHRCPKHWDDADVFNPERWPLDGPNPNETNQKFSYLPFGGGPRKCVGDMFATFETVVATAMLVKRFDFQMA
PGAPPVEMTTGATIHTTKGLNMTVTRRIKPPVIPNLEMKIVSDPEGSTSSTASVAVSTASIASGEGQQVEVS
TSQV

SEQ ID NO: 36: soybean CYP97A Glycine max:

GKGLIPADGEIWRVRRRAIVPALHQKYVAAMIGLFGQAADRLCQKLDAAASDGEDVEMESLFSRLTLDIIGK

AVFNYDFDSLSNDTGIVEAVYTVLREAEDRSVAPIPVWEIPIWKDISPRLRKVNAALKFINDTLDDLIAICK

RMVDEEELQFHEEYMNEQDPSILHFLLASGDDVSSKQLRDDLMTMLIAGHETSAAVLTWTFYLLSKEPRVMS

KLQEEVDSVLGDQYPTIEDMKKLKYTTRVINESLRLYPQPPVLIRRSLEDDVLGEYPIKRGEDIFISVWNLH

RSPKLWDDADKFKPERWALDGPSPNETNQNFKYLPFGGGPRKCVGDLFASYETVVALAMLMRRFNFQIAVGA

PPVEMTTGATIHTTQGLKMTVTHRIKPPIVPSLQMSTLEVDPSISLSDQDEVSQKGEVYQAQAQS

SEQ ID NO: 37: wheat CYP97A Triticum aestivum:
GCRLPQAVGSLASVAGEAFFLPLYDLFLTYGGVFRLNFGPKSFLIVSDPDVAKHILRDNSKAYSKGILAEIL
EFVMGTGLIPADGEVWRVRRRAIVPALHQKYVTAMIGLFGNASDRLCQKLDKAASDGEDVEMESLFSRLTLD
VIGKAVFNYDFDSLSYDNGIVEAVYVTLREAEMRSTSPIPTWEIPIWKDISPRQCPKHWDDADVFNPERWPL
DGPNPNETNQKFSYLPFGGGPRKCVGDMFATFETVVATAMLVKRFDFQMAPGAPPVEMTTGATIHTTKGLNM
TVTRRIKPPVIPNLEMKIVSDSEGSTSSTASVAVSTASIASGEGQQVEVSTSQV

Figure 26b

SEQ ID NO: 38: tomato CYP97A Lycopersicon esculentum:

QFPTHHYSKSRLTLSPKFKGSVSNFTIRCSNSNGKQPESVDEGVKKVEKLLDEKRRAELSARIASGEFTVEQ
SGFPSLLKNGLSKLGVPKEFLEFFSRRTGNYPRIPEAKGSISAIRDEPFFMPLYELYLTYGGIFRLIFGPKS
FLIVSDPSIAKHILKDNSKAYSKGILAEILDFVMGKGLIPADGEIWRVRRRAIVPALHQKYVAAMIGLFGKA
TDRLCKKLDVAATDGEDVEMESLFSRLTLDIIGKAVFNYDFDSLTVDTGIVEAVYTVLREAEDRSVAPIPVW
ELPIWKDISPKLKKVNAALKLINDTLDDLIAICKRMVDEEELQFHEEYMNEKDPSILHFLLASGDEVSSKQL
RDDLMTMLIAGHETSAAVLTWTFYLLSKEPSVMAKLQDEVDSVLGDRLPTIEDLKKLRYTTRVINESLRLYP
QPPVLIRRSIEEDVVGGYPIKRGEDIFISVWNLHRCPNHWEEADRFNPERWPLDGPNPNETNQNFSYLPFGG
GPRKCVGDMFATFENLVAVAMLVQRFDFQMALGAPPVKMTTGATIHTTEGLKMTVTRRSRPPIVPNLEMATL
EVD

SEQ ID NO: 39: green alga CYP97A3 Chlamydomonas reinhardtii: ARRRAVVPALHRKYVMSMVDMFGDCAAHGASATLDKYAASGTSLDMENFFSRLGLDIIGKAVFNYDFDSLAH DDPVIQAVYTLLREAEHRSTAPIAYWNIPGIQFVVPRQKRCQEALVLVNECLDGLIDKCKKLVEEEDAVFGE EFLSERDPSILHFLLASGDEISSKQLRDDLMTMLIAGHETTAAVLTWTLYLLSQHPEAAAAIRKEVDELLGD RKPGVEDLRALKMTTRVINEAMRLYPQPPVLIRRALQDDHFDQFTVPAGSDLFISVWNLHRSPKLWDEPDKF KPERFGPLDSPIPNEVTENFAYLPFGGGRRKCIGDQFALFEAVVALAMLMRRYEFNLDESKGTVGMTTGATI HTTNGLN

Figure 27a

SEQ ID NO: 40: CYP97A Arabidopsis thaliana:

tactgttaaaccagtaacgtactctcggagatcgaactttgtagttttctcgtcgagttctaatggacgag atcctttagaggagaattcagtacctaatggtgtgaaaagcttggagaagcttcaagaagagaagcgtcgt gctqagttatctgctaggattgcttctggagctttcactgtacggaaatctagttttccatctacagtgaa gaatggtttatctaagattggaataccaagcaatgttcttgatttcatgtttgattggactggttctgacc aagactaccccaaggttcctgaggctaaaggctcgattcaggcggtccggaacgaagctttcttcatccct ttqtatqaqcttttccttacttatqqtqqaattttcaqqttqacctttqqqcctaaqtcattcttqatcqt qtcqqatccttctattqctaaacatatattqaaqqacaatqcaaaaqcttactccaaqqqqattttaqctq aaattctagattttgtgatgggaaaaggactcattcctgctgatggggagatatggcgtagacgaaggcgt gccattgttcctgcattgcatcaaaagtatgtagcagctatgattagtttattcggagaagcttcagatag gctttgtcagaagcttgatgctgctgcattgaaaggggaagaagtagagatggaatcactcttctctcgtt tgacacttgatattattggcaaggcggttttcaattacgactttgactcccttactaatgataccggtgtg atcgaggcagtgtacactgttctaagagaagctgaagacagaagtgtttcacctattcctgtttgggacat acccatttggaaagatatttccccacgtcagaggaaagttgctacttccttgaaattaatcaatgacacac ttgatgatttgattgcaacatgcaaqagaatggtagaagaagaggagttgcagtttcacgaggagtatatg aacgaaagagatcctagcatccttcactttcttttagcttcaggagatgatgtctctagtaagcagcttcg tgatgacttgatgacaatgcttatagccggacatgaaacatcggcggcagtattaacatggaccttttacc acaaccaccagtactgatccgtcgttctatagataatgatatacttggagagtatccgataaaaaggggag aggatatetteateteggtttggaatetaeategaagteetetgeattgggatgatgeagagaagtteaat tggaggaccgcggaaatgtataggcgacatgtttgcttcctttgagaatgtggtagcaatcgcaatgctta ttcgaagatttaactttcagattgcaccaggagctcctccggtgaaaatgactacaggagctacaatacac accacagaaggattgaaattgacagtaacaaagaggacaaaacctctggacataccatccgtaccgatact caaacaagctcagatgaagaagcaaaaatcttgtgtttagaacagcaaatgttgaattgttggaacatgacc aatgctttctgattatttatctgcactgtaaaatgcagacaagtaaaatgagaagatttattattctttgg

Figure 27b

SEQ ID NO: 41: CYP97A Arabidopsis thaliana:

atggctatggcctttcctctttcttatactccgacgattactgttaaaccagtaacgtactctcggagatcg aactttgtagttttctcgtcgagttctaatggacgagatcctttagaggagaattcagtacctaatggtgtg aaaaqcttqqaqaaqcttcaaqaaqaqaaqcgtcgtgctgagttatctgctaggattgcttctggagctttc actgtacggaaatctagttttccatctacagtgaagaatggtttatctaagattggaataccaagcaatgtt $\verb|cttgatttcatgtttgattggactggttctgaccaagactaccccaaggttcctgaggctaaaggctcgatt|\\$ ttgacctttgggcctaagtcattcttgatcgtgtcggatccttctattgctaaacatatattgaaggacaat gcaaaagcttactccaaggggattttagctgaaattctagattttgtgatgggaaaaggactcattcctgct gatggggagatatggcgtagacgaaggcgtgccattgttcctgcattgcatcaaaagtatgtagcagctatg attagtttattcggagaagcttcagataggctttgtcagaagcttgatgctgctgcattgaaaggggaagaa gtagagatggaatcactcttctctcgtttgacacttgatattattggcaaggcggttttcaattacgacttt gactcccttactaatgataccggtgtgatcgaggcagtgtacactgttctaagagaagctgaagacagaagt $\tt gtttcacctattcctgtttgggacatacccatttggaaagatatttccccacgtcagaggaaagttgctact$ qatqtctctaqtaaqcaqcttcqtgatgacttqatqacaatgcttataqccggacatgaaacatcggcggca qtattaacatqqaccttttaccttttaacaacqqaaccaaqtqtaqttqccaaacttcaaqaagaggttgat tctgtaattggagatagattcccaaccatacaagatatgaaaaagctgaaatacactactcgagtcatgaat gagtcattgagattatatccacaaccaccagtactgatccgtcgttctatagataatgatatacttggagag tatccgataaaaaggggagggatatcttcatctcggtttggaatctacatcgaagtcctctgcattgggat agttacttacctttcggtggaggaccgcggaaatgtataggcgacatgtttgcttcctttgagaatgtggta gcaatcgcaatgcttattcgaagatttaactttcagattgcaccaggagctcctccggtgaaaatgactaca ggagctacaatacaccaccacagaaggattgaaattgacagtaacaaagaggacaaaacctctggacatacca

SEQ ID NO: 42: rice CYP97A Oryza sativa:

atggeggetacetectetgeggeegeegetgeteeaceteegtgeegettaeteggeteeggteaggeacae ctgcgccttcctccttctgctgctgctgctgcttcagctcgtcgccgcctgctcctccgctgcgccgcc tcgggcggcaacgggaaaggcggtggtggcgacggctccggctccgacccggttcttgaggagcggcggcgg cqqcgccaggctqagctggcgcgcgcattgcgtccggcgagttcaccgcccaaggccccgcgtcagtgctc ctctctctttttcaggtggattgctcccctcgcggtggggcttgccaagctcggcccaccgggggagctcgc egcegegetgeteaceaaggtegeeggtggeggaceggagatacegcaggeggtggggtetatgagtge ggtgacagggcaggctttcttcatcccgctctatgatctcttccttacctatggcggcatctttcgcctcaa tttcggccctaaggtgatgcacaatcagaccaatttgctctccaactcggcaactcccaattttgtgttatt attgatggcctaaactttgttcttttcttgttttccccagtctttcctcattgtctctgatccagctatagc taagcacatcctgagggacaactccaaggcttattccaaggtttttgtgttgtcaatttttggatgtagacgtg tttagagtttgtgatgggtacgggtttgatccctgctgatggggagatttggcgtgttaggaggcgccat tgtaccagcaatgcaccagaaggttctacatcatttctgtaccaggtttagcatgatttgatcttcgggttg tgattgaactgatctgaatttcgctttgcagtacgttaccgcaatgataagtctcttcggatatgcttcaga tcggctctgccagaagttggacaaggcagcaacggatgggaggatgtggagatggaatctttgttctctcg actaacactggatgtcattgggaaggcagtcttcaattatgatttcgactcattgtcttacgataatggaat ttcttaaaattttacctttttttggattgatcaggcagtgtatgtgacactgcgagaagcagaaatgcg gagcacttctcctataccaacttgggaaatacccatatggaaagatatttccccgcggcagaagaaggtcaa tgaagctcttgcgctgataaataagactcttgatgaactaattgacatctgcaaggtgaacttcttttctta tgttctgaccttattatttattttttaaaaaaatcaaggcttttagattggctgctgttactcttgcagaga cttttggcatctggagatgatgtatggtgtacctgcagtttaaaatattatagatctccaaacattctggtc ccgtagctgttcccaaaagaaaatatgttcagcaactcaaatcattgatgtattaatgtagcaatatgtaat gatgaatattgtatacgtcaaaccactcatgttttacctttcttggcattggtaacttgcagtatccaaatg taatggccaaactccaagatgaggtaaattcgcttttacatttaggattgttatttttagaggcacgtgctt ctacatcttacaagttgcaaatgacttgtttcactcacttatggacaggctgatactgttctaggtgaccgt $\verb|ttacca| acaattg | aggatg tg | agaattg | aggta | actacta | aggta | attacca| aggta | aggta |$ agactctatccacagccaccagttttaattcgtcgctctattgaggaggatatgctgggagggtacccaatt ttgttacaattggaataggaaaacagaaagatcatgcctagtatcacaccagtaaagttctggtgaaactga gtctaagggcctctttatttaggcttaagtttattggtttagtattttaagtcagtttttttggcttatatg atttataagccaatggatttaaagtcctaagtttaatggtggagtcatacctctatctcacataagccaaaa aaccttttccaacctagcttttgtcttaatagtgtaacagttcgctgagccttaatagtgtaatagtggttt ttataagcctaaacaaagaggcctaagatatgtgcaagtataagttatctaaccaatctttttttagagaa tatcttcccaatcttgtgtgtatatatttttgtcttctgcttgtataccatttctggtcgtagctctagagta tttaattgtctgaattgttcctttttaaaaatttcagtacaaatttgaacttcacctataattctgataaga tatttttttccctttgtttcccaggggagaagacattttcatatccgtgtggaacctacatcattgcccaaag ${\tt caaaaatttcaggttccatctctattaatgctatgaaatgcattagctctttatttggatgcaccttatcact}$ ctaattcccccattttattagtttctgctttctactacaaaaatcagtagacatttgattatgctcggtata gttgtgttcttgtttgacgagaagtttgcgtttttacatctactaacactaagttatttggtcatgcgatgc acctgttgtaattattctagagataacaaaaacaaaactctagctgattttcgttttcttttgatgcaa atcatcaaattttcttcatgtgattcgtattaatttagtgctaatgctggcatgtgcatagccacaatctta gaaagtgcataaggcagtcacaacaacataaagaaaaatgggacctttttcttttgaagcagtaaagatgag ataactgtgatgacttgattcctaaatttatggttttgggagctaaaccacagtttatgacaatcatgttaa aatgatattcatatggctataagcaacatgtgccaaaatgctattgtactttcaaactgatagtgcttgaaa gtacttttcaactttaccacatcgccagaactgttaactggttacgcagaaacagtacttggaatagtaatt ttgtataaactgatagtatgtcgtatgagttccttacctgtagcaagtatgtcaacagcagaaccctttgta tggatcaacaacatcagaggtgctttgaacaaaatccttacttgattagaagacgtaacaattcgtgccat

Figure 27d

cctgagttaatgaagacttccgactggactacaaacttcttccgtgttgcagtcttgcagggagttgatgca qcaaqtctcctaggcatcacactgaaacttaaaaatggtaatctgcatttacttggcacatgacatgtccca tttcagttggcatgtcaaactttttaagctccaattttaggctgtggtagctttcattctgtgtattgcggc tagcatctgttagctgtcactgcctcactggctaatttaatatttgatgaatctacatagctaaatggga ctcacgtgttctgttgaattctcagttacttgccatttggtggcggaccaaggaaatgtgtaggtgacatgt ttgccactttcgaggtaatttgttttagtttttgaaggatttcttcttttaatttcaaaatgtcattttaag gaaacatagcaaacttatgtatggtccagtcttactgaaccttgttgccttgagccttctgttgtctaccat aaggacattatateteatgeeatgataaataatgtagtacaataaetattgageatgeaagatteeaaetet aataacatggatatgccggaacttgtatgcagactgtggtggcaactgcaatgcttgtcaggcgctttgatt agataatcactgtgaagtatcaatatgataggttgagatgacaactggagcaacgattcacacaactgaggg gttgaaaatgactgttactcggaggacaaagccacctgtaatcccaaacctagagatgaaagtcatttctga ttcaccagaaaacatgagtactactacatcaatgcccgtttctgctgctagtattgcttcaggagaagatca acaagggcaagtctcagcaactcgaatctga

Figure 27e

SEQ ID NO: 43: rice CYP97A Oryza sativa:

 $\verb|atggcggctacctcctctgcggccgccgctgctccacctccgtgccgcttactcggctccggtcaggcacac| \\$ ctgcgccttcctccttctgctgctgctgctgcttcagctcgtcgccgcctgctcctccgctgctgccgcc tcgggcggcaacgggaaaggcggtggtggcgacggctccggctccgacccggttcttgaggagcggcggcgg cggcgccaggctgagctggcgcgcgcattgcgtccggcgagttcaccgcccaaggccccgcgtggattgct cccctcgcggtgggcttgccaagctcggcccaccgggggagctcgccgccgcgctgctcaccaaggtcgcc ccgctctatgatctcttccttacctatggcggcatctttcgcctcaatttcggccctaagtctttcctcatt gtctctgatccagctatagctaagcacatcctgagggacaactccaaggcttattccaagggtattctggca gaaattttagagtttgtgatgggtacgggtttgatccctgctgatggggagatttggcgtgttaggaggcgc gccattgtaccagcaatgcaccagaagtacgttaccgcaatgataagtctcttcggatatgcttcagatcgg ctctgccagaagttggacaaggcagcaacggatgggaggatgtggagatggaatctttgttctctcgacta acactggatgtcattgggaaggcagtcttcaattatgatttcgactcattgtcttacgataatggaatagtt gaggcagtgtatgtgacactgcgagaagcagaaatgcggagcacttctcctataccaacttgggaaataccc ctgatgacaatgctcattgctggccatgagacctctgcagcagtcttgacatggacattttatcttctatct aagtatccaaatgtaatggccaaactccaagatgaggctgatactgttctaggtgaccgtttaccaacaatt gaggatgtgaagaattgaagtatactactagagtaattaacgaatcattgagactctatccacagccacca gttttaattcgtcgctctattgaggaggatatgctgggagggtacccaattggccggggagaagacattttc atatccgtgtggaacctacatcattgcccaaagcattgggatggtgcagatgtttttaatccagaaagatgg cctttggatggaccaaatccaaatgaaacaaaccaaaatttcagttacttgccatttggtggcggaccaagg aaatgtgtaggtgacatgtttgccactttcgagactgtggtggcaactgcaatgcttgtcaggcgctttgat tttcaaatggctccaggagctcctccggttgagatgacaactggagcaacgattcacacaactgaggggttg aaaatgactgttactcggaggacaaagccacctgtaatcccaaacctagagatgaaagtcatttctgattca ccagaaaacatgagtactactacatcaatgcccgtttctgctgctagtattgcttcaggagaagatcaacaa gggcaagtctcagcaactcgaatctga

SEQ ID NO: 44: CYP97A barley Hordeum vulgare:

teggeacgagggeaggeegtegggtegettggetteegteggegggaggeettetteetgeegetetaegae ctcttcctcacctacggeggegtcttccgcctcaacttcgggcccaagtctttcctcatcgtctctgatccg gatgtagctaagcatatcctcagggacaactcaaaggcttattccaagggtatccttgcggaaatactggag tttgtgatgggcacaggtctgatcccggctgatggggaggtctggcgtgttcgacggcgtgccattgtacca gcattgcatcagaagtacgtgacagcgatgataggtctcttttggaaacgcttcagaccggctctgccagaag atcgggaaggcggtgttcaattatgattttgattcattatcttacgataatggaatagttgaggctgtgta tgtaacactgcgggaagcagaaatgcggagtacatctcctattccaacatgggaaatacccatatggaaag cccaacgtaatgtccaagctccaagctgaggctgatgctgttctaggagatggtctgccaacaattgatga tgtgaagaaactgaagtatactactcgagttattaatgaatctttgagactatacccacagccgccagttt taattegeegeteeettgaggatgatatgetaggagagtaeeegateggeaagggagaagatatttttatat ccatctggaaccttcatcgctgcccaaagcattgggatgacgcggatgttttcaatccggaaaggtggcctt tggacggaccgaatccaaatgagacaaaccaaaaattcagttacttgccatttggtggcggaccaaggaaat gtgtaggtgatatgtttgctacttttgagactgtggtagcaacagcaatgcttgtcaagcgatttgattttc agatggctccaggagcacctccggtcgagatgacaaccggagcaacgattcacacaactaagggactgaaca tgactgttactcggaggataaagccacctgtaattccaaacttagagatgaaaatcgtttccgatccagaag gaagcacaagttctactgcgtcagtggctgtttctactgctagtattgcatccggagaaggtcaacaagtgg aggtgtcgacaagtcaagtgtga

Figure 27f

SEQ ID NO: 45: soybean CYP97A Glycine max:

qqqaaaqqqcttatcccaqctqatqqtqaaatatqqcqaqttaqacqtcqtqctataqtcccaqcattqcac cagaagtatgtagcagctatgattggccttttcggacaagctgcagataggctctgccagaagctagatgct gctgcatccgatggagaagatgttgagatggaatcacttttctctcgattgaccttggacatcattggcaag gcagtattcaattatgattttgatagtttatcaaatgacactggtatagttgaggctgtttatactgtactg agagaagcagaagatcgaagtgttgctccaattccagtctgggagatcccaatatggaaagacatatcacca cgtctaaggaaggttaatgcagctctcaaattcatcaatgatacgcttgatgatctgatagcaatatgcaag ttcttgttggcgtcaggagatgatgtgtcaagtaagcaacttcgtgatgacttaatgaccatgctcattgct ggacatgaaacatcagctgctgttttaacttggaccttttatcttctatccaaggagccaagagtcatgtcc aagctccaagaagaggttgactctgtacttggagatcaatatccaactatagaagacatgaagaaactcaaa tatacaacccgagtgatcaatgagtcattgaggctttacccacaaccacctgtgttaattcgccgctctctt gaggatgatgttcttggagagtaccctataaagagaggtgaagatatctttatatctgtatggaacctgcat cgcagtccaaaactatgggatgatgctgacaagtttaaacctgaaagatgggcattagatggaccaagtcct aatgagacaaatcaaaacttcaaatatcttccgtttggtggcggaccacggaaatgtgtaggtgatttgttt gcttcatacgagacggtagtagcactcgcaatgcttatgagacgattcaactttcaaatagcagttggagct ccaccggttgagatgactactggagctacaattcatacaacacaagggttgaagatgactgtaactcacaga caagatgaagtaagtcagaaaggcgaagtttaccaggctcaggctcagtcctaa

SEQ ID NO: 46: wheat CYP97A Triticum aestivum:

Figure 27g

tomato CYP97A Lycopersicon esculentum: SEQ ID NO: 47: caatttccaacaccattactctaaatctagactcactctctcacctaaattcaagggtagtgtatcaaat tttacaattaqqtqttctaattctaatgqaaaacagcctgagtcggtagatgaaggagtcaaaaaggtggaa aaqcttttaqatqaqaaaaqqcqaqctqaattatctqctcgtattgcttcaggcgaatttactgttgaacaa ${\tt tctggcttcccgtcattgctcaaaatggtttgtctaaattgggtgtaccaaaggaatttcttgagttcttc}$ tctcgacgaacgggcaattatcctcgcattccagaggcaaaaggatccatcagtgctattcgggatgagcca $\verb|ttcttcatgccgctttatgagctttaccttacttatggcggaattttccggttgatttttggtcccaagtct|\\$ tttttaataqtttctgatccatcaatagccaaacacatactgaaagataattctaaggcttattctaagggt atcctagctgaaatattggactttgtgatgggaaagggacttatacctgcagatggagaaattttggcgcgtc aggcggcgtgccattgtaccagcattgcaccaaaagtacgtagcagctatgattggcttatttggaaaaagca accgataggttgtgcaaaaagcttgatgttgctgcaactgatggagaagatgtagagatggaatcacttttc tcccgtctaacattggacatcattggcaaagctgtatttaattatgattttgactctttaactgtagatactggtatcgtggaggctgtatatacagtacttagagaagcagaagatcgtagtgttgcaccaattccagtttgg gagttgcctatctggaaagatatctctccgaagctaaaaaaggttaatgcagctctcaagttgattaatgac acattggatgatctgattgctatatgtaagaggatggtagacgaagaagagttgcagtttcacgaggaatac $\verb|cgtgatgacctcatgacaatgcttatagcgggacatgaaacatctgcagcagtgctcacatggaccttttat|\\$ ctqttqtccaaqqaacctaqtqtcatgqccaagcttcaagatgaggtcgattcagttctaggggataggtta ccaaccattqaaqatctaaaqaaactcaqatacacaactcgtgtgattaatgagtctttaagactatatcca cagccaccagtcttgattcgtcgttctattgaagaggacgtagttggaggttacccgattaaaaggggtgaa qacatttttcatttctgtttggaacttgcatcgatgcccgaatcattgggaagaagccgatagattcaatcct ggaccaagaaaatgtgtgggagacatgtttgccacatttgagaatttagtagcagttgcaatgcttgttcaa cgatttgattttcaaatggctcttggagctcctcctgttaaaatgacaactggggctaccatccacaccaca gaaggattaaaaatgactgtaacacgaagatcaagacctccaatagttcccaacttggagatggcaacatta gaagtagat

CYP97A like gene Chlamydomonas reinhardtii: SEQ ID NO: 48: gcgcgccgacgcgcagtggtgccagccctgcaccgcaagtacgtgatgtcgatggtggacatgttcggcgac tgcgcggcgcacggcgcgccacactagacaagtatgccgcctcaggcaccagcctggacatggaaaac $\verb|ttcttcagccggctgggtctggacatcatcggcaaggccgtgttcaactacgacttcgactcgctggcgcac| \\$ qacqaccccqtcatccaqqccqtqtacacqttqctqcgcgaaqcggagcaccgctccacagcgcccatcgcc tactggaacattcccggcatccagtttgtggtgccgcggcagaagcgctgccaggaggcgctggtgctggta aatgagtgcctggacggcctcatcgacaagtgcaagaagctggtcgaggaggaggacgcggtgtttggggag qaqttccttaqcqaqcqcqacccctccatcctqcacttcctcctcqcqtctqqaqacqaqatttcctcqaag cagttgcgcgatgacctgatgactatgctgattgcggggcacgagaccaccgccgccgtgctgacgtggacg ctgtacctgctgtcccaacaccccgaggcggcagcggccatccgcaaggaggtagacgagctccttggggac cgcaagcccggggtggaagacctcagagcgctcaagatgacgactcgcgtcatcaacgaggcgatgcggctc tacccacagccgccagtactcattcgccgcgcgctgcaggacgaccacttcgaccagttcacggtgccggcc $\tt ggcagcgacctgttcatcagcgtgtggaacttgcaccgcagccctaagctgtgggacgagcccgacaagttc$ aagccggagcgcttcggaccgctggacagccccatccccaacgaggtgactgaaaacttcgcctacctgccc tttqqcqqtqqccqccqcaaqtqcattqqcqaccaqttcgctttgttcqaqqcggttgttgcgctggccatg $\verb|ctgatgcggcgatacgagttcaacctggacgagtccaaggggacagtggccatgacaacaggtgccaccatc|\\$ cacaccaccaacggtctaaac

Figure 28

SEQ ID NO: 49: CYP97B3 Arabidopsis thaliana:

MAFPAAATYPTHFQGGALHLGRTDHCLFGFYPQTISSVNSRRASVSIKCQSTEPKTNGNILDNASNLLTNF
LSGGSLGSMPTAEGSVSDLFGKPLFLSLYDWFLEHGGIYKLAFGPKAFVVISDPIIARHVLRENAFSYDKG
VLAEILEPIMGKGLIPADLDTWKLRRRAITPAFHKLYLEAMVKVFSDCSEKMILKSEKLIREKETSSGEDT
IELDLEAEFSSLALDIIGLSVFNYDFGSVTKESPVIKAVYGTLFEAEHRSTFYFPYWNFPPARWIVPRQRK
FQSDLKIINDCLDGLIQNAKETRQETDVEKLQERDYTNLKDASLLRFLVDMRGVDIDDRQLRDDLMTMLIA
GHETTAAVLTWAVFLLSQNPEKIRKAQAEIDAVLGQGPPTYESMKKLEYIRLIVVEVLRLFPQPPLLIRRT
LKPETLPGGHKGEKEGHKVPKGTDIFISVYNLHRSPYFWDNPHDFEPERFLRTKESNGIEGWAGFDPSRSP
GALYPNEIIADFAFLPFGGGPRKCIGDQFALMESTVALAMLFQKFDVELRGTPESVELVSGATIHAKNGMW
CKLKRRSK

SEQ ID NO: 50: pea CYP97B1 and CYP97A2 Pisum sativum:

MVAAPISTVKLTDANLHTRFHSSSSSTPSTLSLPLSLHFHFSSHSKRFSSIRCQSVNGEKRKQSSRNVFDN
ASNLLTSLLSGANLGSMPIAEGAVTDLFDRPLFFSLYDWFLEHGSVYKLAFGPKAFVVVSDPIVARHILRE
NAFSYDKGVLADILEPIMGKGLIPADLETWKQRRRVIAPGFHTSYLEAMVQLFTSCSERTVLKVNELLEGE
GRDGQKSVELDLEAEFSNLALEIIGLGVFNYDFGSVTNESPVIKAVYGTLFEAEHRSTFYIPYWKFPLARW
IVPRQRKFQDDLKVINTCLDGLIRNAKESRQETDVEKLQQRDYSNLKDASLLRFLVDMRGVDVDDRQLRDD
LMTMLIAGHETTAAVLTWAVFLLAQNPDKMKKAQAEVDLVLGMGKPTFELLKKLEYIRLIVVETLRLYPQP
PLLIRRSLKPDVLPGGHKGDKDGYTIPAGTDVFISVYNLHRSPYFWDRPNDFEPERFLVQNNNEEVEGWAG
FDPSRSPGALYPNEIISDFAFLPFGGGPRKCVGDOFALMESTVALVCCYRISMWN

SEQ ID NO: 51: soybean CYP97B2 Glycine max:

MSVDTSSTLSTVTDANLHSRFHSRLVPFTHHFSLSQPKRISSIRCQSINTDKKKSSRNLLGNASNLLTDLL
SGGSIGSMPIAEGAVSDLLGRPLFFSLYDWFLEHGAVYKLAFGPKAFVVVSDPIVARHILRENAFSYDKGV
LADILEPIMGKGLIPADLDTWKQRRRVIAPAFHNSYLEAMVKIFTTCSERTILKFNKLLEGEGYDGPDSIE
LDLEAEFSSLALDIIGLGVFNYDFGSVTKESPVIKAVYGTLFEAEHRSTFYIPYWKIPLARWIVPRQRKFQ
DDLKVINTCLDGLIRNAKESRQETDVEKLQQRDYLNLKDASLLRFLVDMRGADVDDRQLRDDLMTMLIAGH
ETTAAVLTWAVFLLAQNPSKMKKAQAEVDLVLGTGRPTFESLKELQYIRLIVVEALRLYPQPPLLIRRSLK
SDVLPGGHKGEKDGYAIPAGTDVFISVYNLHRSPYFWDRPDDFEPERFLVQNKNEEIEGWAGLDPSRSPGA
LYPNEVISDFAFLPFGGGPRKCVGDQFALMESTVALTMLLQNFDVELKGTPESVELVTGATIHTKNGLWCN
LRKRSSLH

SEQ ID NO: 52: rice CYP97B4 Oryza sativa:

MAAAAAAAVPCVPFLCPPPPPLVSPRLRRGHVRLRLRPPRSSGGGFTGGGGAGGDEPPITTSWVSPDWLTA
LSRSVATRLGGGDDSGIPVASAKLDDVRDLLGGALFLPLFKWFREEGPVYRLAAGPRDLVVVSDPAVARHV
LRGYGSRYEKGLVAEVSEFLFGSGFAIAEGALWTVRRRSVVPSLHKRFLSVMVDRVFCKCAERLVEKLETS
ALSGKPVNMEARFSQMTLDVIGLSLFNYNFDSLTSDSPVIDAVYTALKEAELRSTDLLPYWKIDLLCKIVP
RQIKAEKAVNIIRNTVEDLITKCKKIVDAENEQIEGEEYVNEADPSILRFLLASREEVTSVQLRDDLLSML
VAGHETTGSVLTWTIYLLSKDPAALRRAQAEVDRVLQGRLPRYEDLKELKYLMRCINESMRLYPHPPVLIR
RAIVDDVLPGNYKIKAGQDIMISVYNIHRSPEVWDRADDFIPERFDLEGPVPNETNTEYRFIPFSGGPRKC
VGDQFALLEAIVALAVVLQKMDFTIELVPDQKINMTTGATIHTTNGLYMNVVNIGVQVDEARKHGYNSFIV
YGYTLYAYISPRIWSAMPVL

Figure 29a

SEQ ID NO: 53: CYP97B3 in Arabidopsis thaliana:

atqqcttttcctgccgctgctacttatcccacccatttccaaggcggcgctcttcatctgggtaggaccgat cattgcctcttcggtttctaccctcaaaccatttcctctgtgaattctcggagagcttctgtttccatcaag tgccaatctacggagccaaagacgaatggtaacatattggacaatgcgagcaaccttttgacaaatttttta agtggtggaagtttggggtcaatgcctactgctgaaggctctgtctctgatttgttttggaaagcctctcttt ttatctctttacgactggttcttggagcatggaggaatttataaacttgcgtttggtccaaaagcctttgtt gtcatctcagatcccattattgcaaggcatgtcctccgggaaaatgctttttcttatgacaagggagttctt gctgagatcttagagccgattatgggaaaagggttaataccggctgatctagatacgtggaagttaagaaga agagctatcactcccgcattccataaattgtatctagaggccatggtcaaagtatttagtgactgttcggag gatctggaagcagaattctcgagtctggctcttgatattataggtcttagcgtgttcaactacgatttttggc tctgtcacaaaagagtcccctgtgatcaaggcagtttatggaactcttttcgaggcagagcatcggtctact ttctacttcccttattggaactttcctccagctagatggatagttccgaggcaacgaaagttccaaagcgat ctgaagattataaacgattgccttgatggcctcattcaaaatgctaaagagacaagacagcaggaaacagat gttgagaagctccaggaaagggactacactaatctcaaggatgcaagtcttttgcggttcttagtcgatatg cgcggtgttgacattgatgaccggcagctgagggatgacttgatgactatgctaattgctggtcatgagaca acagcagcagtacttacttgggctgttttccttctgtcacaaaatcctgaaaaaattaggaaagctcaagct gagattgatgctgtgcttggtcaaggtccacccacttatgaatcaatgaaaaagctcgagtacatacgactg ategttgtagaagteettegtetettteeteageeacetttgeteateagaegeaeteteaaaceagaaaca ttacccggtggacacaaaggggaaaaagaaggtcataaagttccaaaagggactgatatcttcatttctgta tataateteeatagateteeataettttgggataateeecaegattttgageetgagaggtttttaagaaea aaggagagcaatggaattgaaggatgggctggctttgatccatctcgtagccccggggcactatatccgaat gagataatagcagactttgcattcttaccatttggtggaggaccaagaaaatgcattggagaccagtttgca ctaatggaatcgaccgtcgcactagctatgttgtttcagaaattcgatgtggagctgcgtggaacgccagaa tctgttgaactcgtgagcggcgcaacgattcatgccaaaaatgggatgtggtgcaaactaaagagaagatca aagtga

SEQ ID NO: 54: CYP97B1 and CYP97A2 for Pisum sativum: catcacttaccactaactgaaacttgcaagcaccattctcaacttaacaccgtcgtcaccgccatggttgc cgcccctatctcaaccgtcaaacttaccgatgccaatcttcacaccagatttcattcctcttcttcta caccatccaccctcagtcttccactctctcttcattttcacttttcttctccactccaaacgcttttcttct atcagatgtcaatcggttaatggtgaaaagcgaaaacaaagtagtagaaatgtgtttgacaatgctagcaa $\verb|cctccttacaagcttgttaagtggtgcaaatttagggtccatgcccatagctgaaggtgccgtcacagatc|\\$ tgtttgaccggccgctgtttttctcactatatgattggttcttagagcatggttctgtgtataaactggcg tttggaccgaaagcatttgttgttgtatcagatcccattgttgcaagacatattctgcgagaaaatgcatt ttettatgacaagggagtaettgetgatateetagaaceaattatgggaaaaggaeteataeetgeagaee ttgagacatggaagcaaaggagaagagtgattgctccgggtttccatacctcatacttggaagctatggta caactattcacttcatgttcagaaagaactgtgttaaaggtcaatgagcttcttgaaggagaggggcgtga tggacagaagtcagttgaattggaccttgaggcagaattttcaaatttggctcttgagattattgggctag gtgtgttcaactatgactttggttctgtcaccaatgaatctcccgttattaaggctgtctatggcactctt tttgaagccgaacatagatccactttctatattccatattggaaatttccattagcaaggtggattgtgcc caggcaaaggaagtttcaggatgaccttaaagtcattaatacttgtcttgatggacttatcagaaatgcaa aagagagcaggcaggaaacagatgttgagaaactgcagcaaagggattactcaaatttgaaggatgcaagt cttctgcgtttcctagttgatatgcggggagttgatgttgatgatcgtcagttgagggatgatttaatgac aatgcttattgctggtcatgagacgacggctgcagttcttacatgggcagttttcctgctagctcaaaatc ctgacaaaatgaagaaggctcaagcagaggtagattttggtgctggggatggggaagccaactttttgaattg cttaaaaagttggagtacattaggttaattgttgtggagactcttcgattatatccacaaccacctctgct gattagacgttcactcaaacctgatgttttgccaggtggacataaaggtgacaaagatggttatacaattc ctgctgggactgatgtcttcatttctgtatataatctccatcgatctccatattttttgggaccgccctaat gacttcgagcctgaacgatttctagtgcaaaacaataatgaagaagttgaagggtgggctggttttgaccc atctcgaagtcctggagccttgtatccaaacgagattatatcagatttttgcattcttgccttttggtggtg gaccacgaaaatgcgttggagaccaatttgctctcatggaatccactgtagcgctagtatgctgctacaga atttcgatgtggaactgaaggggacccctgaatcggttgaactagttactggggcaactatccataccaaa aatggattgtggtgcaatttgaggaagagatctagtttacattgacatgttaactgcaacatttttcttat gcagaatgatgtacaaaatatttatcatttaaaatgacattaacattgaatagtgtctaatacagctaaag

Figure 29b

SEQ ID NO: 55: soybean CYP97B2 Glycine:

atgagtqtcqacacttcctccaccctctccaccqtcaccqatqccaatcttcactccagatttcattctcg tcttqttccattcactcatttctcactttctcaacccaaacqqatttcttcaatcaqatqccaatcaa ttaataccqataaqaaqaaatcaaqtaqaaatctqctqqqcaatqcaaqtaacctcctcacqqacttatta agtggtggaagtatagggtctatgcccatagctgaaggtgcagtctcagatctgcttggtcgacctctctt $\verb|tttctcactgtatgattggttcttggagcatggtggtgtataaacttgcctttggaccaaaagcatttg|$ ttgttgtatcagatcccatagttgctagacatattctgcgagaaaatgcattttcttatgacaagggagta cttqctqatatccttqaaccaataatqqqcaaaqqactcataccagcagaccttgatacttggaagcaaag qaqaaqaqtcattqctccqqctttccataactcatacttqqaaqctatqqttaaaatattcacaacttgtt cagaaagaacaatattgaagtttaataagcttcttgaaggagagggttatgatggacctgactcaattgaa ttqqatcttqaqqcaqaqttttctaqtttgqctcttgatattattgggcttggtgttcaactatgactt tggttctgtcaccaaagaatctccagttattaaggcagtctatggcactctttttgaagctgaacacagat gatgacctaaaggtcatcaatacttgtcttgatggacttatcagaaatgcaaaagagagcagacaggaaaac agatgttgagaaattgcagcagagggattacttaaatttgaaggatgcaagtcttctgcgtttcctggttg atatgcggggagctgatgttgatgatcgtcagttgagggatgatttaatgacaatgcttattgccggtcat qaaacaacqqctqcaqttcttacttqqqcaqttttcctcctaqctcaaaatcctaqcaaaatqaaqaaggc $\verb|tcaagcagaggtagatttggtgctgggtacggggaggccaacttttgaatcacttaaggaattgcagtaca|\\$ $\verb|ttagattgattgttgtggaggctcttcgtttatacccccaaccacctttgctgattagacgttcactcaaa|$ $\verb|tctgatgttttaccaggtgggcacaaaggtgaaaaagatggttatgcaattcctgctgggactgatgtctt|\\$ catttctqtatataatctccataqatctccatatttttqqqaccqccctqatqacttcgaaccagagagat ttcttgtgcaaaacaagaatgaagaaattgaaggatgggtcttgatccatctcgaagtcccggagcc ttgtatccgaacgaggttatatcggattttgcattcttaccttttggtggcggaccacgaaaatgtgttgg qqaccaatttqctctqatggagtccactgtagcgttgactatgctgctccagaattttgacgtggaactaa a agggacccctgaatcggtggaactagttactggggcaactattcataccaaaaatggaatgtggtgcagattgaagaagagatctaatttacgttga

Figure 30

SEQ ID NO: 56: novel cytochrome P450 monooxygenase diatom CYP97B Skeletonema costatum:

MASYESDLLSTWDEDPSLQKGFDWEIEKLRRYFAGLRQTPDGRWVRKSTLFEFLVTNSPSKVVGVGPDGER
YESPPKPVNIFDVGVLVGKNTLTWLGFGPNLGMAAVPDAVIQKYEGSFFTFIKGALGGDLQTLAGGPLFLL
LAKYYTDHGPIFNLSFGPKSFLVISDPVMARHILRDSSPEQYCKGMLAEILEPIMGDGLIPADPKIWKVRR
RAVVPGFHKKWLNSMIGLFGDCGDRLVDDLEKRSTSDKPVIDMEERFCSVTLDIIGKAVFNYDFGSVTKES
PIVKAVYRVLREAEHRSSSFIPYWNLPYAEKWMVGQVEFRKDMGMLDDILAKLINRAVETRQEATVEELEE
RETSDDPSLLRFLVDMRGEDLTSKVLRDDLMTMLIAGHETTAAMLTWTMFGLVSNDPGMMKEIQAEVRTVM
GNKSRPDYDDVVAMKKLRYALIEALRLYPEPPVLIRRARQEDTLPPGGTGLSGGVKVLRGTDIFISTWNLH
RAPEYWENADKYDPTRWERPFKNPGVKGWNGYDPEKQSSQSLYPNEITSDYAFLPFGAGKRKCIGDQFAML
EASVTLSMIMNKFDFTLVGTPEDVGMKTGATIHTMNGLNMMVSPRSETNPIPGTNEWWTKQHLMRGLSSTG
RPYTSDEDAAWTTSANGMRP

Figure 31

SEQ ID NO: 57: novel cytochrome P450 monooxygenase diatom CYP97B Skeletonema costatum:

atggcctcctacgagagtgatctgctctcaacatgggatgaagatccatcgctgcaaaaggggtttgactg qqaqattqaaaaqctccqtcqqtactttqccqqactgcgtcaaacaccagacgggcgatgggtgcgcaagt cgacactgtttgagtttcttgtgacaaactctccaagtaaagtagttggggtaggtccggatggggaacgg tatqaaaqccctccqaaaccaqtcaatatcttcqatqtgggagtgttagtcggtaagaatacactcacttg qttqqqatttqqaccqaatttqqqtatqqccqcqqtacccgatgcagtcattcaaaagtatgagggtagct tcttcacctttatcaagggagcattggggggtgatttgcaaactttggcgggtggtcctttgttcttattg cttqccaagtattatacggatcatggacccattttcaacttgagtttttggaccaaagagctttttggtgat ttcqqatcctqttatgqcgaggcatattttgagggatagttcaccggagcagtattgtaagggaatgcttg cggagattttggaaccgatcatgggtgatggattgattcctgcagatccaaagatttggaaggttcgtcga agagctgtcgtacctggtttccacaaaaagtggctgaacagcatgattggtttgttcggagactgtggtga tcgtctcgttgacgatctagaaaagcgttctacttcagataaacctgtaattgacatggaagaacgattct gttccgtcacactcgatatcatcggtaaggcagtattcaactatgattttggatcagtgacaaaggaatca $\verb|cctattgtaaaggcagtatacagagtgttacgtgaggcggagcacagatcatcttcgttcatcccctactg|$ gaacttgccttatgctgagaaatggatggtaggacaggttgaattccgcaaagatatgggaatgcttgacg atatettqqcaaaactqatcaatcqtqctqttqaqactaqqcaaqaaqetactqtcqaaqaqttqqaaqaq aqaqaaacaaqcqatqatccqaqtctcttaaqqttcctagttgatatgaggggagaagatttaacgagtaa aqtqttqaqaqatqatttqatqacaatqcttattqcaqqacatgaaacaacagcggcaatgctgacgtgga caatqtttqqqctaqtaaqcaacqatcctqqcatqatqaaqqaaatccaqqcaqaagttcgaactgtcatg acttcgattatatcccgagccacccgtgttgattcgcagggcaaggcaagaggacactcttccaccaggtg qtacqqqtctttctqqaqqtqtcaaaqtattqcqtqqaacaqatatctttatttctacttqgaaccttcac cqcqctccaqaatactqqqaqaatqcaqacaaatatgaccctactcgatqqqaqcqtccgttcaaaaaaccc aggtgttaagggttggaatggatatgatccggaaaaacaatcatctcaatcactttatcctaacgagataa cgtcagactatgctttcctttccttttggtgctgggaagagaaaatgtatcgggggatcagtttgctatgctc qaqqcttcqqttacactatcqatqattatgaataaatttgacttcacqttggtcggtacccctgaagatgt cggcatgaagaccggagcaactattcataccatgaatgggctcaacatgatggtcagccctcgatcagaga caaacccgattccagggacaaatgagtggtggacgaaacaacatctaatgagaggtttgagttctactgga agaccatacacttccgatgaagatgccgcgtggacgacatccgctaatggcatgagaccgtga

Fig. 32
SEQ ID NO: 58: single knockout mutant CYP97A3 Arabidopsis thaliana (SALK_116660):

